Amendments to the Specification

Please replace paragraph 268 (page 63 of the PCT publication) with the following amended paragraph:

[0268] The ORF of bnKCP1 gene codes for a 215 amino acid polypeptide product of polypeptide with several functional motifs (FIG. 11). Based on a search of protein localization sites using PSORT program (see URL:~~http://~~psort.nibb.acjp; Nakai and Kanehisa, 1992), bnKCP1 appears to be is a nuclear protein containing a pat7 nuclear localization signal (NLS) PLNKKRR (SEQ ID NO: 62; FIG. 10A, residues 127-133). Three acidic motifs (I, II and III) aid a serine-rich (S-rich) region (residues 34-58) may function in transcription activation by bnKCP1 (Johnson et al., 1993). The charged motif GKSKS (residues 88-143), which is conserved in all four protein orthologs (FIG. 10A), is rich in basic residues and encompasses the NLS. This suggests that this domain serve the may function of a DNA-binding motif (FIG. 11). In addition, bnKCP1 is extremely hydrophilic (FIG. 11) suggesting bnKCP1 is an active element in the nuclear matrix.

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Please replace paragraph 294 (page ~~78~~ 71 of the PCT publication) with the following amended paragraph:

[0294] BnIAA1 and BnIAA12 are clones ML2798 and ML4744, which are homologs of Arabidopsis IAA1 and IAA12, respectively, and were identified in a database of Brassica napus ESTs that were generated at the Saskatoon Research Centre of Agriculture and Agri-Food Canada (see URL:~~www~~.brassica.ca).

Please replace paragraph 303 (page 75 of the PCT publication) with the following amended paragraph:

[0303] Comparison of the deduced BnSCL1 amino acid sequence to the NCBI (see URL:~~http://www~~.ncbi.nln.nih.gov) and TAIR (see URL:arabidopsis.org) databases results in a list of proteins with considerable similarity (FIG. 21). According to the NTI computer program